## **Amendments to the Claims:**

This listing of claims will replace all prior versions and listings of claims in the application.

## **Listing of Claims:**

1. (currently amended) A method for screening for compounds that affect uncoupling, comprising:

contacting a mammalian cell or tissue sample with a candidate compound; analyzing expression of a polypeptide having at least 9095% sequence identity to a polypeptide encoded by SEQ ID NO:1 or 2 wherein the polypeptide having at least 95% sequence identity has uncoupling activity; and

analyzing mitochondrial membrane potential,

wherein a change in expression of the polypeptide <u>having at least 95% sequence identity</u> indicates that the compound affects uncoupling.

2-27. (canceled)

28. (previously presented) The method of claim 1, wherein the mammalian cell or tissue sample is a human cell or tissue sample.

29-33. (canceled)

- 34. (previously amended) The method of claim 1, wherein the analyzing of expression of the polypeptide comprises analyzing expression of a polypeptide having at least 95% amino acid sequence identity to the polypeptide encoded by SEQ ID NO:1.
- 35. (previously amended) The method of claim 1, wherein the analyzing of expression of the polypeptide comprises analyzing expression of a polypeptide having at least 95% amino acid sequence identity to the polypeptide encoded by SEQ ID NO:2.

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- 36. (previously amended) The method of claim 28, wherein the analyzing of expression of the polypeptide comprises analyzing expression of a polypeptide having at least 95% amino acid sequence identity to the polypeptide encoded by SEQ ID NO:1.
- 37. (previously amended) The method of claim 28, wherein the analyzing of expression of the polypeptide comprises analyzing expression of a polypeptide having at least 95% amino acid sequence identity to the polypeptide encoded by SEQ ID NO:2.
- 38. (previously presented) The method of claim 1, wherein the candidate compound is a member selected from the group consisting of a small molecule, a polynucleotide, a modified polynucleotide, a polypeptide, an antibody, an antibody fragment and a modified antibody.
- 39. (currently amended) The method of claim 1 A method for screening for compounds that affect uncoupling, comprising:

contacting a mammalian cell or tissue sample with a candidate compound;
analyzing expression of a polypeptide, wherein the polypeptide is encoded by
SEQ ID NO:1; and
analyzing mitochondrial membrane potential,
wherein a change in expression of the polypeptide indicates that the compound
affects uncoupling.

40. (currently amended) The method of claim 1 A method for screening for compounds that affect uncoupling, comprising:

contacting a mammalian cell or tissue sample with a candidate compound; analyzing expression of a polypeptide, wherein the polypeptide is encoded by SEQ ID NO:2; and analyzing mitochondrial membrane potential, wherein a change in expression of the polypeptide indicates that the compound affects uncoupling.

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41. (previously presented) A method for screening for compounds that affect uncoupling, comprising:

contacting a mammalian cell or tissue sample with a candidate compound; and analyzing expression of a polypeptide encoded by SEQ ID NO:1 or 2, wherein a change in expression of the polypeptide indicates that the compound affects uncoupling.

- 42. (canceled)
- 43. (previously presented) The method of claim 41, further comprising analyzing mitochondrial membrane potential in the sample.
- 44. (currently amended) A method for screening for compounds that affect uncoupling, comprising:

contacting a mammalian cell or tissue sample with a candidate compound suspected of affecting uncoupling; and

analyzing expression of a polypeptide having at least 9095% sequence identity to a polypeptide encoded by SEQ ID NO:1 or 2 wherein the polypeptide having at least 95% sequence identity has uncoupling activity,

wherein a change in expression of the polypeptide <u>having at least 95% sequence identity</u> indicates that the compound affects uncoupling.

- 45. (canceled)
- 46. (previously presented) The method of claim 44, wherein the mammalian cell or tissue sample is a human cell or tissue sample.
  - 47-51. (canceled)

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- 52. (previously amended) The method of claim 44, wherein the analyzing of expression of the polypeptide comprises analyzing expression of a polypeptide having at least 95% amino acid sequence identity to the polypeptide encoded by SEQ ID NO:1.
- 53. (previously amended) The method of claim 44, wherein the analyzing of expression of the polypeptide comprises analyzing expression of a polypeptide having at least 95% amino acid sequence identity to the polypeptide encoded by SEQ ID NO:2.
- 54. (previously amended) The method of claim 46, wherein the analyzing of expression of the polypeptide comprises analyzing expression of a polypeptide having at least 95% amino acid sequence identity to the polypeptide encoded by SEQ ID NO:1.
- 55. (previously presented) The method of claim 46, wherein the analyzing of expression of the polypeptide comprises analyzing expression of a polypeptide having at least 95% amino acid sequence identity to the polypeptide encoded by SEQ ID NO:2.
- 56. (previously presented) The method of claim 44, wherein the candidate compound is a member selected from the group consisting of a small molecule, a polynucleotide, a modified polynucleotide, a polypeptide, an antibody, an antibody fragment and a modified antibody.
- 57. (currently amended) The method of claim 44, A method for screening for compounds that affect uncoupling, comprising:

contacting a mammalian cell or tissue sample with a candidate compound

suspected of affecting uncoupling; and

analyzing expression of a polypeptide, wherein the polypeptide is encoded by

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SEQ ID NO:1; and

wherein a change in expression of the polypeptide indicates that the compound affects uncoupling.

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58. (currently amended) The method of claim 44, A method for screening for compounds that affect uncoupling, comprising:

contacting a mammalian cell or tissue sample with a candidate compound suspected of affecting uncoupling; and analyzing expression of a polypeptide, wherein the polypeptide is encoded by SEQ ID NO:2; and wherein a change in expression of the polypeptide indicates that the compound affects uncoupling.

## 59.-73. (cancelled)

- 74. (new) The method of claim 41, wherein the analyzing of expression of the polypeptide comprises analyzing expression of a polypeptide encoded by SEQ ID NO:1.
- 75. (new) The method of claim 41, wherein the analyzing of expression of the polypeptide comprises analyzing expression of a polypeptide encoded by SEQ ID NO:2.
- 76. (new) The method of claim 41, wherein the candidate compound is a member selected from the group consisting of a small molecule, a polynucleotide, a modified polynucleotide, a polypeptide, an antibody, an antibody fragment and a modified antibody.